



ENTERED

OIPE

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/766,511B

DATE: 05/16/2002 P.6  
TIME: 17:03:46

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\05162002\I766511B.raw

3 <110> APPLICANT: MCCARTHY, Sean A  
 4 FRASER, Christopher C  
 5 SHARP, John D  
 6 BARNES, Thomas S  
 7 KIRST, Susan J  
 8 MYERS, Paul S  
 9 WRIGHTON, Nicholas  
 10 GOODEARL, Andrew  
 11 HOLTZMAN, Douglas A  
 12 KHODADOUST, Mehran M  
 14 <120> TITLE OF INVENTION: NOVEL GENES ENCODING PROTEINS HAVING PROGNOSTIC, DIAGNOSTIC,  
 PREVENTIVE,  
 15 THERAPEUTIC, AND OTHER USES  
 17 <130> FILE REFERENCE: 10147-65  
 19 <140> CURRENT APPLICATION NUMBER: 09/766,511B  
 C--> 20 <141> CURRENT FILING DATE: 2002-05-07  
 22 <150> PRIOR APPLICATION NUMBER: US 09/578,063  
 23 <151> PRIOR FILING DATE: 2000-05-24  
 25 <150> PRIOR APPLICATION NUMBER: US 09/333,159  
 26 <151> PRIOR FILING DATE: 1999-06-14  
 28 <150> PRIOR APPLICATION NUMBER: US 09/596,194  
 29 <151> PRIOR FILING DATE: 2000-06-16  
 31 <150> PRIOR APPLICATION NUMBER: US 09/342,364  
 32 <151> PRIOR FILING DATE: 1999-06-29  
 34 <150> PRIOR APPLICATION NUMBER: US 09/608,452  
 35 <151> PRIOR FILING DATE: 2000-06-30  
 37 <150> PRIOR APPLICATION NUMBER: US 09/393,996  
 38 <151> PRIOR FILING DATE: 1999-09-10  
 40 <150> PRIOR APPLICATION NUMBER: US 09/345,680  
 41 <151> PRIOR FILING DATE: 1999-06-30  
 43 <160> NUMBER OF SEQ ID NOS: 85  
 45 <170> SOFTWARE: PatentIn version 3.1  
 47 <210> SEQ ID NO: 1  
 48 <211> LENGTH: 2964  
 49 <212> TYPE: DNA  
 50 <213> ORGANISM: Homo sapiens  
 52 <400> SEQUENCE: 1  
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 57 ggacgcgagg agccatgagg cgccagcctg cgaagggtggc ggcgctgctg ctcgggctgc 180  
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69 cgccctatta cactgaccca ggaggaccgg ggatgaaccc tgcgggaat tccatggcaa 540
71 tggctttcca ggccccccc aactcacccc aggggagtggt ggccctgccc cccctccag 600
73 cctactgcaa cagcctccg ccccgtagc aacaggtagt gaaggccaag tagtggggtg 660
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83 cccgaggggt gacgtcctta cgggtggcgtg accagatcta caggagagag actgagagga 960
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155 <211> LENGTH: 516
156 <212> TYPE: DNA
157 <213> ORGANISM: Homo sapiens
159 <400> SEQUENCE: 2
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162 gaagccaaaa agcattgctg gtatttcgaa ggactctatc caacctatta tatatgccgc 120
164 tcctacgagg actgctgtgg ctccagggtgc tgtgtgcggg ccctctccat acagaggctg 180
166 tggactttct ggttccttct gatgatgggc gtgcttttct gctgcgggagc cggcttcttc 240
168 atccggaggc gcatgtaccc ccgcccgtg atcgaggagc cagccttcaa tgtgtcctac 300
170 accaggcagc ccccaaatcc cggcccagga gccagcagc cggggccgcc ctattacact 360
172 gacccaggag gaccggggat gaacctgtc gggaattcca tggcaatggc tttccaggtc 420
174 ccacccaact caccccaggg gagtgtggc tgcccgcgcc ctccagccta ctgcaacacg 480
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182 &lt;213&gt; ORGANISM: Homo sapiens

184 &lt;400&gt; SEQUENCE: 3

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187 1 5 10 15
190 Leu Glu Cys Thr Glu Ala Lys Lys His Cys Trp Tyr Phe Glu Gly Leu
191 20 25 30
194 Tyr Pro Thr Tyr Tyr Ile Cys Arg Ser Tyr Glu Asp Cys Cys Gly Ser
195 35 40 45
198 Arg Cys Cys Val Arg Ala Leu Ser Ile Gln Arg Leu Trp Tyr Phe Trp
199 50 55 60
202 Phe Leu Leu Met Met Gly Val Leu Phe Cys Cys Gly Ala Gly Phe Phe
203 65 70 75 80
206 Ile Arg Arg Arg Met Tyr Pro Pro Pro Leu Ile Glu Glu Pro Ala Phe
207 85 90 95
210 Asn Val Ser Tyr Thr Arg Gln Pro Pro Asn Pro Gly Pro Gly Ala Gln
211 100 105 110
214 Gln Pro Gly Pro Pro Tyr Tyr Thr Asp Pro Gly Gly Pro Gly Met Asn
215 115 120 125
218 Pro Val Gly Asn Ser Met Ala Met Ala Phe Gln Val Pro Pro Asn Ser
219 130 135 140
222 Pro Gln Gly Ser Val Ala Cys Pro Pro Pro Pro Ala Tyr Cys Asn Thr
223 145 150 155 160
226 Pro Pro Pro Pro Tyr Glu Gln Val Val Lys Ala Lys
227 165 170

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230 &lt;210&gt; SEQ ID NO: 4

231 &lt;211&gt; LENGTH: 22

232 &lt;212&gt; TYPE: PRT

233 &lt;213&gt; ORGANISM: Homo sapiens

235 &lt;400&gt; SEQUENCE: 4

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237 Met Arg Arg Gln Pro Ala Lys Val Ala Ala Leu Leu Leu Gly Leu Leu
238 1 5 10 15
241 Leu Glu Cys Thr Glu Ala
242 20

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245 &lt;210&gt; SEQ ID NO: 5

246 &lt;211&gt; LENGTH: 150

247 &lt;212&gt; TYPE: PRT

248 &lt;213&gt; ORGANISM: Homo sapiens

250 &lt;400&gt; SEQUENCE: 5

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252 Lys Lys His Cys Trp Tyr Phe Glu Gly Leu Tyr Pro Thr Tyr Tyr Ile
253 1 5 10 15
256 Cys Arg Ser Tyr Glu Asp Cys Cys Gly Ser Arg Cys Cys Val Arg Ala
257 20 25 30
260 Leu Ser Ile Gln Arg Leu Trp Tyr Phe Trp Phe Leu Leu Met Met Gly
261 35 40 45
264 Val Leu Phe Cys Cys Gly Ala Gly Phe Phe Ile Arg Arg Arg Met Tyr
265 50 55 60
268 Pro Pro Pro Leu Ile Glu Glu Pro Ala Phe Asn Val Ser Tyr Thr Arg
269 65 70 75 80
272 Gln Pro Pro Asn Pro Gly Pro Gly Ala Gln Gln Pro Gly Pro Pro Tyr
273 85 90 95
276 Tyr Thr Asp Pro Gly Gly Pro Gly Met Asn Pro Val Gly Asn Ser Met
277 100 105 110
280 Ala Met Ala Phe Gln Val Pro Pro Asn Ser Pro Gln Gly Ser Val Ala
281 115 120 125
284 Cys Pro Pro Pro Pro Ala Tyr Cys Asn Thr Pro Pro Pro Tyr Glu
285 130 135 140
288 Gln Val Val Lys Ala Lys
289 145 150
292 <210> SEQ ID NO: 6
293 <211> LENGTH: 38
294 <212> TYPE: PRT
295 <213> ORGANISM: Homo sapiens
297 <400> SEQUENCE: 6
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300 1 5 10 15
303 Cys Arg Ser Tyr Glu Asp Cys Cys Gly Ser Arg Cys Cys Val Arg Ala
304 20 25 30
307 Leu Ser Ile Gln Arg Leu
308 35
311 <210> SEQ ID NO: 7
312 <211> LENGTH: 21
313 <212> TYPE: PRT
314 <213> ORGANISM: Homo sapiens
316 <400> SEQUENCE: 7
318 Trp Tyr Phe Trp Phe Leu Leu Met Met Gly Val Leu Phe Cys Cys Gly
319 1 5 10 15
322 Ala Gly Phe Phe Ile
323 20
326 <210> SEQ ID NO: 8
327 <211> LENGTH: 91
328 <212> TYPE: PRT
329 <213> ORGANISM: Homo sapiens
331 <400> SEQUENCE: 8
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334 1 5 10 15
337 Val Ser Tyr Thr Arg Gln Pro Pro Asn Pro Gly Pro Gly Ala Gln Gln
338 20 25 30

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TIME: 17:03:47

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\05162002\I766511B.raw

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342          35          40          45
345 Val Gly Asn Ser Met Ala Met Ala Phe Gln Val Pro Pro Asn Ser Pro
346          50          55          60
349 Gln Gly Ser Val Ala Cys Pro Pro Pro Ala Tyr Cys Asn Thr Pro
350 65          70          75          80
353 Pro Pro Pro Tyr Glu Gln Val Val Lys Ala Lys
354          85          90

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357 &lt;210&gt; SEQ ID NO: 9

358 &lt;211&gt; LENGTH: 0

359 &lt;212&gt; TYPE: DNA

360 &lt;213&gt; ORGANISM: Homo sapiens

362 &lt;400&gt; SEQUENCE: 9

W--&gt; 363 000

365 &lt;210&gt; SEQ ID NO: 10

366 &lt;211&gt; LENGTH: 0

367 &lt;212&gt; TYPE: DNA

368 &lt;213&gt; ORGANISM: Homo sapiens

370 &lt;400&gt; SEQUENCE: 10

W--&gt; 371 000

373 &lt;210&gt; SEQ ID NO: 11

374 &lt;211&gt; LENGTH: 2915

375 &lt;212&gt; TYPE: DNA

376 &lt;213&gt; ORGANISM: Mus sp.

378 &lt;400&gt; SEQUENCE: 11

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379 gtcgacccac gcgtccggcc gcgcgtcctt ctgccggctt cagctcgtat ccccgagatc      60
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383 ggtcctcgct ggagccatgg gccgcgggct cggcagggtg gcggcgctgc tgctcgggct      180
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387 atactatata tgccgttcct atgaagactg ctgtggctcc aggtgctgtg tgagggccct      300
389 ttccatacag aggtgtgtgt atttttggtt cctgctgatg atgggtgtgc tgttctgtg      360
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411 gtgttagggg gcagataaag tggtcaggct gagataagac tcacatgatg cagtagttgg      1020
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421 aggggaaatt ctctaaatgg agacattgct ttttatgaat catcgtctgg cttttctttt      1320
423 agtgcattga ttgaagttag ggtgtccttt gagatcagat ggggagagtg aactctgcgg      1380
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RAW SEQUENCE LISTING ERROR SUMMARY      DATE: 05/16/2002  
PATENT APPLICATION:    US/09/766,511B      TIME: 17:03:48

Input Set : A:\PTO.VSK.txt  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:61; N Pos. 788

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 14

**VERIFICATION SUMMARY**

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Input Set : A:\PTO.VSK.txt

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L:20 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:363 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (9) SEQUENCE:  
L:371 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (10) SEQUENCE:  
L:560 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (14) SEQUENCE:  
L:642 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (19) SEQUENCE:  
L:3193 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:780